

**LISTING OF CLAIMS:** This listing of claims replaces all prior versions and listings of claims in the instant patent application.

1. (previously presented) A compound 12 to 50 nucleobases in length targeted to a region comprising nucleotide 901 to 950 of the nucleic acid molecule encoding diacylglycerol acyltransferase 2 in SEQ ID NO: 4, wherein said compound is at least 80% complementary to said nucleic acid molecule encoding diacylglycerol acyltransferase 2, and wherein said compound comprises at least an 8 nucleobase portion of SEQ ID NO: 35, 36, 37 or 38.
2. (canceled)
3. (currently amended) The compound of claim 1, wherein said compound is comprising 15 to 30 nucleobases in length.
4. (original) The compound of claim 1 comprising an oligonucleotide.
5. (currently amended) The compound of claim 4 comprising an ~~antisense~~ antisense oligonucleotide.
6. (original) The compound of claim 4 comprising a DNA oligonucleotide.
7. (original) The compound of claim 4 comprising a RNA oligonucleotide.
8. (original) The compound of claim 4 comprising a chimeric oligonucleotide.
9. (original) The compound of claim 4 wherein at least a portion of said compound hybridizes with RNA to form an oligonucleotide-RNA duplex.
10. (canceled).
11. (original) The compound of claim 1 having at least 90% complementarity with said nucleic acid molecule encoding diacylglycerol acyltransferase 2.

12. (currently amended) The compound of claim 1 having at least 95% complementarity with said nucleic acid molecule encoding diacylglycerol ~~acyltransferase~~ acyltransferase 2.

13. (currently amended) The compound of claim 1 having 100% complementarity with said nucleic acid molecule encoding diacylglycerol ~~acyltransferase~~ acyltransferase 2.

14. (currently amended) The compound of claim 1 ~~havig~~ having at least one ~~modified~~ modified internucleoside linkage, sugar moiety, or nucleobase.

15. (original) The compound of claim 1 having at least one 2'-O-methoxyethyl sugar moiety.

16. (original) The compound of claim 1 having at least one phosphorothioate internucleoside linkage.

17. (original) The compound of claim 1 having at least one 5-methylcytosine.

18.-21. (canceled)

22. (original) A kit or assay device comprising the compound of claim 1.

23.-43. (canceled)

44. (original) The compound of claim 1, wherein said compound comprises an antisense nucleic acid molecule that is specifically hybridizable with a coding region of the diacylglycerol acyltransferase 2 (SEQ ID NO: 4).

45. - 57. (canceled)

58. (previously presented) The compound of claim 1, wherein said compound is 20 nucleobases in length.

59. (currently amended) The compound of claim 13 having at least one modified ~~intenucleoside~~ internucleoside linkage, sugar moiety, or nucleobase.

60. (previously presented) A compound 20 nucleobases in length targeted to a nucleic acid molecule encoding diacylglycerol acyltransferase 2 (SEQ ID NO: 4), wherein said compound has the nucleobase sequence of SEQ ID NO: 35.

61. (New) The compound of claim 60, wherein said compound is an antisense oligonucleotide comprising:

- a gap segment consisting of linked deoxynucleosides;

- a 5' wing segment consisting of linked nucleosides;

- a 3' wing segment consisting of linked nucleosides;

wherein the gap segment is positioned between the 5' wing segment and the 3' wing segment and wherein each nucleoside of each wing segment comprises a modified sugar.

62. (New) The compound of claim 61, wherein the antisense oligonucleotide comprises:

- a gap segment consisting of ten linked deoxynucleosides;

- a 5' wing segment consisting of five linked nucleosides;

- a 3' wing segment consisting of five linked nucleosides;

wherein the gap segment is positioned between the 5' wing segment and the 3' wing segment, wherein each nucleoside of each wing segment comprises a 2'-O-methoxyethyl sugar; and wherein each internucleoside linkage is a phosphorothioate linkage.

63. (New) The compound of claim 62, wherein each cytosine is a 5-methylcytosine.